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Minimum
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Database
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Run on:
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                                                                                                                                                                                                                                                                                                                                                          Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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1644.5
1638.5
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125.5
125.5
125.5
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125.5
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146
138.5
138.5
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138.5
138.5
137
137
137
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1: /cgn2_6/ptodata/2,
2: /cgn2_6/ptodata/2,
3: /cgn2_6/ptodata/2,
4: /cgn2_6/ptodata/2,
5: /cgn2_6/ptodata/2,
6: /cgn2_6/ptodata/2,
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                   US-09-587-111-5
4004
1 MTSPSSSPVERLETLDGGQE......EDEDGASEENYVPVQLLQSN 764
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Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           193259 seqs, 20144635 residues
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6A_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/BCTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/backfiles1.pep:*
                                                                                                                                                                                                                                                                                                        Length
DB
                                      US-09-082-059-2
US-09-172-977-3
US-09-172-096-15
US-09-065-474-139
US-09-065-474-33
US-08-847-429A-33
US-08-623-679-9
US-08-931-485-23
US-08-847-429A-23
US-09-065-474-29A-23
US-09-065-474-29A-23
US-09-065-474-29A-23
US-09-065-474-29A-23
US-09-065-474-29A-23
US-09-065-474-29A-23
US-09-065-474-29A-28
US-09-065-474-29A-28
US-08-897-340-32
US-08-897-340-32
US-08-897-340-32
US-08-699-103B-25
US-08-699-103B-25
                   US-08-537-210A-4
US-09-113-825-4
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4, Appli
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15, Appl
13, Appl
133, Appl
33, Appl
33, Appl
33, Appl
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28, Appl
32, Appl
32, Appl
33, Appl
32, Appl
33, Appl
32, Appl
33, Appl
33, Appl
                 Appl:
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US-09-197-636-2
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Result

No

	119.5	120.5	120.5	120.5	120.5	120.5	120.5				120.5			120.5	120.5	122	123.5	123.5
	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.0	3.1	3.1
	357	741	741	741	741	741	741	741	741	741	741	741	302	302	302	741	191	191
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	US-08-031-538-4	PCT-US95-02058-4	PCT-US95-02058-2	US-08-479-895-2	US-08-701-005A-2	US-08-487-797-4	US-08-487-797-2	US-08-434-998-4	US-08-434-998-2	US-08-436-771-4	US-08-436-771-2	US-08-462-481-2	US-09-065-474-38	US-08-847-429A-38	US-09-031-485-38	US-08-943-956A-2	US-09-065-474-20	US-08-847-429A-20
•	Sequence 4, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 2, Appli	•	Sequence 38, Appl	Sequence 38, Appl	Sequence 2, Appli	Sequence 20, Appl	Sequence 20, Appl

ALIGNMENTS

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; MOLECULE TYPE: US-09-197-636-2
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                                                                                                                                                                                       APPLICATION NUMBER: US/09/197,636
FILING DATE: 23-NOV-1998
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 9805137.8
FILING DATE: 12-MAR-1998
APPLICATION NUMBER: UK 9815791.0
FILING DATE: 21-UIL-1998
APPLICATION NUMBER: UK 9819278.4
FILING DATE: 03-SEP-1998
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
REFERENCE/DOCKET NUMBER: 23,031
REFERENCE/DOCKET NUMBER: GP-30075
TELECHONE: 601-407-0700
                                                                                                                                 TELEFAX: 610-407-0701
TELEX: 846169
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: DUCKWORTH, DAVID
APPLICANT: HAYES, PHILIP
APPLICANT: MEADOWS, HELEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TITLE OF INVENTION: NOVEL COMPOUNDS NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 839 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT:
                                     TYPE: amino acid STRANDEDNESS: sir TOPOLOGY: linear
                                                                                                                                                        TELEPHONE: 6U1-70
TELEPHONE: 610-407-0701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Ratner & Prestia
STREET: P.O. Box 980
CITY: Valley Forge
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US-09-197-636-8
                                                                                          APPLICANT: DUCKWORTH, DAVID
APPLICANT: HAYES, PHILIP
APPLICANT: MEADOWS, HELEN
APPLICANT: DAVIS, JOHN
APPLICANT: DAVIS, JOHN
TITLE OF INVENTION: NOVEL COMPOUNDS
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                Sequence 8, Application US/09197636 Patent No. 6239267
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Best Local Similarity
                                                                                                                                                                                                                                                             GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   633
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    173 TIPLLLEIARQTDSLKELVNASYTDSYYKGQTALHIAIERRNMALVTLLVENGADVQAAA
                                     CITY: Valley Forge
STATE: PA
                    COUNTRY:
                                                                              ADDRESSEE: Ratner & Prestia STREET: P.O. Box 980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   74 FDRDRLFNAVSRGVPEDLAGLPEYLSKTSKYLTDSEYTEGSTGKTCLMKAVLNLKDGVNA 133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CILPLLQIDRDSGNPQPLVNAQCTDDYYRGHSALHIAIEKRSLQCVKLLVENGANVHARA 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ESKNIWKLQRAITILDTEKSFLKCMRKAFRSGKLLQVGYTPDGKDDYRWCFRVDEVNWTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MVVLEPLNKLLQAKWDLLIPK-FFLNFLCNLIYMFIFTAVAYHQPTLKKQAAPHLKAE-V
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VEVADNTADNTKFVTSMYNEILILGAKLHPTLKLEELTNKKGMTPLALAAGTGKIGVLAY
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  19482-0980
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48.5%; Pred. No. 9.16
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TELEX: 846169
INFORMATION FOR SEO ID NO: 8:
SEQUENCE CHARACTERISTICS:
'ENGTH: 839 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 41.1%; Score 1644.5; DB 4. Best Local Similarity 48.5%; Pred. No. 1.1e-146; Matches 344; Conservative 121; Mismatches 200;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION NUMBER: UK
APPLICATION NUMBER: 12-MAR-190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
TELEPHONE: 601-407-0700
TELEFAX: 610-407-0701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Ver:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/197,636
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: UK 9815791.0 FILING DATE: 21-UUL-1998 APPLICATION NUMBER: UK 9819278.4 FILING DATE: 03-SEP-1998 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
590
                                       548
                                                                                                                                                                                                                                                                                                                     353
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
SOFTWARE: FastSEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Prestia, Paul F
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                       MLLVEPLNRLLQDKWDRFVKRIFYFNFLVYCLYMIIFTMAAYYRPV----DGLPPFKMEKT
GFSTAVVTLIEDGKNDSLPSESTSHRWRGPACRPPDSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CGRFFQKGQG-TCFYFGELPLSLAACTKQWDVVSYLLENPHQPASLQATDSQGNTVLHAL 252
                                   GFAVALVSLSQEA----
                                                                          YESHLKEYVASMVESLALGWTNMLYYTRGFQQMGIYAVMIEKMILRDLCREMEVYVVFLF
                                                                                              CFLAIEWYLPLLVSALVLGWLNLLYYTRGFQHTGIYSVMIQKVILRDLLRFLLIYLVFLF
                                                                                                                                                                                             GNSMLLTGHILILLGGIYLLVGQLWYFWRRHVFIWISFIDSYFEILFLFQALLTVVSQVL 487
                                                                                                                                                                                                                                                                            MVVLEPLNKLLQAKWDLLIPK-FFLNFLCNLIYMFIFTAVAYHQPTLKKQAAPHLKAE-V 427
                                                                                                                                                                                                                                                                                                                 YDRRSIFEAVAQNNCQDLESLLLFLQKSKKHLTDNEFKDPETGKTCLLKAMLNLHDGQNT 172
                                                                                                                                                        GDYFRVTGEILSVLGGVYFFFRGIQYFLQRRPSMKTLFVDSYSEMLFFLQSLFMLATVVL
                                                                                                                                                                                                                                                                                                                                                                                                                      VMISDNSAENIALVTSMYDGLLQAGARLCPTVQLEDIRNLQDLTPLKLAAKEGKIEIFRH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TIPLLLEIARQTDSLKELVNASYTDSYYKGQTALHIAIERRNMALVTLLVENGADVQAAA 232
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                                   -WRPEAPTGPNATESVQPMEGQEDEGNGAQYRGIL 594
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                                                                          Matches
                                                                                        Best Local Similarity
                                                                                                         Query Match
                                                                                                                                                                                                                                                                                TELEX: 846169
INFORMATION FOR SEQ ID NO: 4:
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: UK 980
FILING DATE: 12 - MAR-1998
APPLICATION NUMBER: UK 981
FILING DATE: 21 - UL-1998
APPLICATION NUMBER: UK 981
FILING DATE: 03 - SEP-1998
APPLICATION UMBER: UK 981
FILING DATE: 03 - SEP-1998
ATTORNEY/AGENT INFORMATION:
NAME: Prestia, Paul F
REGISTRATION NUMBER: 23,03
                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 839 amino acid
                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: GP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 601-407-0700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: MEADOWS, HELEN
APPLICANT: DAVIS, JOHN
TITLE OF INVENTION: NOVEL COMPOUNDS
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
     113
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                  74 FDRDRLFNAVSRGVPEDLAGLPEYLSKTSKYLTDSEYTEGSTGKTCLMKAVLNLKDGVNA 133
                                                                                                                                                                                                                                                                                                        TELEPHONE: 6U1 TC.
TETERAX: 610-407-0701
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STREET: P.O. Box 980
CITY: Valley Forge
                                                                                                                                                                                              TOPOLOGY:
                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: si
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version
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YDRRSITEAVAQNNCQDLESLLLFLQKSKKHXTDNEFKDPETGKTCLLKAMLNLHDGQNT 172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DSWSIWKLQKAISVLEMENGYWWC-RKKQRAGVMLTVGTKPDGSPDERWCFRVEEVNWAS
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                                                                      Conservative 121; Mismatches 201;
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                                                                                                                                                                                                               single
                                                                                        40.9%; Score 1638.5; DB 4 48.3%; Pred. No. 4.2e-146;
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                                                                                                         DB 4;
                                                                      Indels
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; FEATURE: -
; OTHER INFORMATION: 929491
US-09-172-977-4
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                                                                                             NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 4
LENGTH: 1839
                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                  APPLICANT: Tang, Y. Tom
APPLICANT: Guegler, Karl J.
APPLICANT: COrley, Neil C.
APPLICANT: Yue, Henry
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILLE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
CURRENT FILING DATE: 1998-10-14
                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/09172977 Patent No. 5989863
                                                  TYPE: PRT
ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GNSMLLTGHILLLGGIYLLVGQLMYFWRRHVFIWISFIDSYFEILFLFQALLTVVSQVL 487
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GDYFRVTGEILSVLGGVYFFFRGIQYFLQRRPSMKTLFVDSYSEMLFFLQSLFMLATVVL
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Query Match 3.8
Best Local Similarity 27.3
Matches 88; Conservative

3.8%;

46;

Score 151; DB 2; Pred. No. 6.9e-05; 6; Mismatches 96;

Length 1839; Indels

92;

Gaps

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RESULT 6
US-09-172-977-3
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US-09-082-059-2
                                                                                           Sequence 3, Application US/09172977 Patent No. 5989863
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/09082059A Patent No. 6225086 GENERAL INFORMATION:
                                                                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 1088
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: 60/047356
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ. ID NOS: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Morrow, Jon S.

APPLICANT: Devarajan, Prasad
TITLE OF INVENTION: No. 6225086el Ankyrin Proteins and a Method for Their Identificat
FILE REFERENCE: 44574-5002-US.
APPLICANT: Corley, Neil C
                          APPLICANT:
                                             APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
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                                                                                                                                                                                                                                   181 ITTK-----KGFTPLHVAAKYGKLEVANLLLQKSAS
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                                                                                                                                                                                                                                                                                                                                                                                                                         RGETALHMAARSGQAEVVRYLVQDGAQVEAKA-----KDDQT------PLHISARLGK 132
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SGLSHLSRKFTEWCYGPVRVSL----YDLASVDSCEENSVLEIIAFHCKSPHRHRMVVLE 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RAGQVEVVRC----LLRNGA-----LVDARAREEQTPLHIASRLGKTEIVQLLLQH-- 519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LQIDRDSGNPQPLVNAQCTDDYYRGHSALHIAIEKRSLQCVKLLVENGANVHARACGRFF 198
                                                                                                                                                                                                                                                                                                                               ADIVQQLLQ---QGASPNAATTSGYTPLH---LSAREGHEDVAAF---
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PLH--VAAKYGSLDVAKLLLQRRAAADSAGKNGLTPL-----HVAAHYDNQKVALLLL 616
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SAENIALVTSMYDGLLQAGARLCPTVQLEDIRNLQDLTPLKLAAKEGKIEIFRHILQREF 318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----ITESGLTPIHVAAFMGHLNIVLLLLQN---GASPDVTNIRGETALHMAA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        QKGQGTCFYFGELPLSLAACTKQWDVVSYLLENPHQPASLQATDSQGNTVLHALVMISDN 258
                        Tang, Y. Tom
Guegler, Karl J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3.7%; Score 150; DB 4; Length 1088; 32.3%; Pred. No. 3.8e-05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US/09/082,059A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51;
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            Query Match
Best Local Similarity
Matches 151; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Yue, Henry
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
CURRENT FILING DATE: 1998-10-14
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SOFTWARE: PERL Program
                                                                                                                                                                                                      SOFTWARE: FastSEQ for Windows Version 3.0 SEQ ID NO 15 ... LENGTH: 1095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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Best Local Similarity
                                                                                                                                                                                                                                                                               EARLIER APPLICATION NUMBER: 60/056,110 EARLIER FILING DATE: 1997-08-20 NUMBER OF SEQ ID NOS: 29
                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Prostate Tumor Polynucleotide TITLE OF INVENTION: Antiqen Compositions FILE REFERENCE: 7636-0015.30
CURRENT APPLICATION NUMBER: US/09/112.096
CURRENT FILING DATE: 1998-07-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Reiner Laus
APPLICANT: Michael H. Shapero
APPLICANT: Larisa Tsavaler
                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 843
TYPE: PRT
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OTHER INFORMATION: 91841966
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Rattus norvegicus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 416 KKQAAPHLKAEVGNSMLLTGHI 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 531 PLH--VAAKYGSLDVAKLLLQRRAAADSAGKNGLTPL-----HVAAHYDNQKVALLLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   392 -----ITESGLTPIPVAAFMGHLNIVLLLLQN---GASPDVTNIRGETALHMAA----- 437
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            348 LLLDK-RANP----NARALN----GFTPLHIACKKNRIKVMELLVKYGAYIQA-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          EKGASPHATAKNGYTPL---HI 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RAGEVEVVRC----LLRNGA-----LVDARAREEQTPLHIASRLGKTEIVQLLLQH--
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                        Conservative
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27.3%; Pred. No. 3.9e-05;
45. Mismatches 97;
                                           3.6%;
                        126;
                   Score 146; DB 4; I
Pred. No. 9.1e-05;
Pred. Mismatches 215;
                                                              Length 1095;
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                        Indels 354;
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RESULT 8
US-09-065-474-139
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               APPLICANT: Tang, Liang
APPLICANT: Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULE
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 171
                                                                                                                                                                                                                                                                                                                                               1042
  CORRESPONDENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       844
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                                                                                                                                                                                                                                                                                                                                            ETLAWE 1047
                                                                                                                                                                                                                                                                                                                                                                                          NWASWE 715
                                                                                                                                                                                                                                                                                                                                                                                                                                        YCSRLNIPFPFIVFAYFYMVVKKCFKCCCKEKNMESSVC--
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SKQWYGEISRDTKNWKIILCLFIIPLVGCGFVSFRKKPVDKHKKLLWYYVAFFTSPFVVF 729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KKQAAPHLKAEVGN-SMLLTGHILILLGGIYLLVGQ-----LWYF-----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CEA-----GSNCLELAVEATDQHFTAQPGVQNFL 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CKSPHRHRMVVLEPLNKLLQAKWDLLIPKFFLNFLCNLIYMFI-----FTAVAYHQPTL 415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TLAALGASKLLKTLAKVKNDINAAGESEELANEYETRAVELFTECYSSDEDLAEQLLVYS 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            VRVSL-----SCEENSVLEIIAFH 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IRN-----SGLSHLSRKFTEWCYGP 335
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SW--NVVFYIAFLLLFAYVLLMDFHSVPHPPELVLYSLVFVLFCDEVRQWY----VNGVN 783
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N------HFSTLVYRNLQIAKNS-YNDALLTFVWKLVANFRRGFR-----KE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SRGVPEDLAG-LPEYLSKTSKYLTDS-----EYTEGSTGKTCLMKAVLNLKDG-----V
ADDRESS
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                                                                     MOLECULES,
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US-09-031-485-33; Sequence 33, Application US/09031485; Patent No. 5824306; GENERAL INFORMATION:
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Best Local
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NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HW-5-C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
                                                                                                                                         APPLICANT: Tang, Liang
APPLICANT: Blebm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            241
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          147 LHVAAHYNNDKVALLLLENGASAHAAAKN-------GYTPLHIAAKKNQMDIAS 193
                                                                                                                                                                                                                                                                                                                                                                                                                 295 IE--HG-ARVSVITRASYTPLHQAAQQGHNSVVRYLLEHGASPNVHTSTGQTPLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      227 YLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQL 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               167 LHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVS 226
                                   STATE: C
                                                                        STREET: 1825 Sharp CITY: Fort Collins
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ADDRESSEE: Heska Corporation
STREET: 1825 Sharp Point Drive
CITY: Fort Collins
                                                                                                            ADDRESSEE: Heska Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: 2. CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     TEWCYGPVRVSLYDLASVDSCEE-----NSVLEIIAFHCKSPHRHRMVVLEPLN 377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -----NGLTPMHLCAQEDRVSVAEELVKENAAIDPKTKAGYTPLHVACHFGQINMVRFL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EDIRNLQDLTPLKLAAKEGKIEIFRHILQREFS--------GLSHLSRKF 328
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                 80525
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Similarity 25.1%; Pred. No. 7.7
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                                                                                        1825 Sharp Point Drive
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MEDIUM TYPE:

E: Floppy disk IBM PC compatible

US/08/847,429A

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                                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 33, Application US/08847429A Patent No. 5827692
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
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                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
                                                                                                                                                                                                                                                                                                                         APPLICANT: Blebm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
TITLE OF INVENTION: USES THEREOF
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS: LENGTH: 1745 amino acids
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         730 IE--HG-ARVSVITRASYTPLHQAAQQGHNSVVRYLLEHGASPNVHTSTGQTPLS 781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        582 LHVAAHYNNDKVALLLLENGASAHAAAKN------GYTPLHIAAKKNQMDIAS 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 167 LHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVS 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     329 TEWCYGPVRVSLYDLASVDSCEE-----NSVLEIIAFHCKSPHRHRMVVLEPLN 377
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COPTWARE: WordPerfect for Windows, Version 7.0
                                                                                                                                                                                                 CITY: Fort Collins
                                                                                                                                                                                                                   ADDRESSEE: Carol Talkington Verser, Ph.D. ADDRESSEE: Heska Corporation STREET: 1825 Sharp Point Drive
                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 970/484-9505
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
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                      WordPerfect for Windows, Version 7.0
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INFORMATION FOR SEQ ID NO:
                            ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
                                                                                                                                                                                                                           ZIP: 80525
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
TITLE OF INVENTION: USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459 REFERENCE/DOCKET NUMBER: HW-5 TELECOMMUNICATION INFORMATION: TELEPHONE: 970/493-7272
                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Tang, Liang APPLICANT: Blehm, E S
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: protein
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              676 -----NGLTPMHLCAQEDRVSVAEELVKENAAIDPKTKAGYTPLHVACHFGQINMVRFL 729
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           629 TLL---HYKANANAESKAGFTPLH--LAAQEGHREMAAL-----LIENGAKVGAQAR- 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     227 YLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQL 286
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       329 TEWCYGPVRVSLYDLASVDSCEE-----NSVLEIIAFHCKSPHRHRMVVLEPLN 377
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                                                                                               APPLICATION NUMBER: FILING DATE: 24-APR CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Sin
                                                                                                                                                                                                                                                                                                                    STATE:
                                                                                                                                                                                                                                                                                                                                          CITY:
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                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Carol Talkington ADDRESSEE: Heska Corporation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/0
FILING DATE: 24-APR-1997
                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH:
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                                                                                                                                                                                                                                                                                                                                      Fort Collins
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59; Conservative
                                                                                                                                                                                                                                                                                                                    Colorado
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                                                                                                                 24-APR-1998
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25.1%; Pred. No. 0.00097;
7ative 33; Mismatches 84;
                                                                                                                                                                                                                                                                                                                                                                                                Talkington Verser,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scot
                                                                                                                                      US/09/065,474
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                  HW-5-C1
                                                                                                                                                                             Version
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RESULT 12
US-08-623-679-9
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TELEX: 200154
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 1533 amin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/08623679 Patent No. 5674739
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 1745 amino aci
                                                                           TELEPHONE: 617/542-8906
TELEXAMINATION: 11/542-8906
TELEXAMINATION: 70/542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Shyjan, Andrew W.
TITLE OF INVENTION: COMPOSITITIE OF INVENTION: DIAGNOSITITITLE OF INVENTION: PROGRESS
                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/412,431
FILING DATE: 29-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Fasse, J. Peter
                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Boston
STATE: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     629 TLL---HYKANANAESKAGFTPLH--LAAQEGHREMAAL-----LIENGAKVGAQAR- 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            582 LHVAAHYNNDKVALLLLENGASAHAAAKN-------GYTPLHIAAKKNQMDIAS 628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             167 LHIAIEKRSLQCVKLLVENGANVHARACGRFFQKGQGTCFYFGELPLSLAACTKQWDVVS 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    730 IE--HG-ARVSVITRASYTPLHQAAQQGHNSVVRYLLEHGASPNVHTSTGQTPLS 781
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     227 YLLENPHQPASLQATDSQGNTVLHALVMISDNSAENIALVTSMYDGLLQAGARLCPTVQL 286
                                                                                                                                                              NAME: Fasse, J. Peter REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 02110-2804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Match 3.5%; Local Similarity 25.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/623,679
                                                                                                                                                                                                                                                                                                                                                     SOFTWARE:
                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -----NGLTPMHLCAQEDRVSVAEELVKENAAIDPKTKAGYTPLHVACHFGQINMVRFL 729
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                                                                                                                                                                                                                                                                                                                                                                       IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                               29-MAR-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PROGRESSION
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                                                                                                                                                07334/004001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    84;
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; TYPE: PRT; ORGANISM: Homo sapiens US-08-933-774-9
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US-08-623-679-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-933-774-9
Query Match
Best Local Similarity
Matches 101; Conserv
                                                                                                                                                                                                          SOFTWARE: FastSEQ SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Shyjan, Andrew W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9, Application US/08933774A Patent No. 6025137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS, PREVENTION TITLE OF INVENTION: AND TREATMENT OF TUMOR PROGRESSION FILE REFERENCE: 07334/004003 CURRENT APPLICATION NUMBER: US/08/933,774A CURRENT FILING DATE: 1997-09-19 EARLIER APPLICATION NUMBER: US 08/623,679 EARLIER FILING DATE: 1996-03-29
                                                                                                                                                                                                                                                  EARLIER APPLICATION NUMBER: US 08/412,431
EARLIER FILING DATE: 1995-03-29
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                  LENGTH: 1533
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         998 VNLLIAVFNNTFFEVKSISNQVWKFQR 1024
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      426 EVGNSMLLTGHILILLGGIYLLV----GQLWYFWRRHVFIWISFIDSYFEI------LF 474
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              262 NIA-----LVTSMYDGLLQ------AGARLCPTVQLEDIRNLQDLTPLKLAAKEG 305
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HKKQRSI--PIGTKICEFYNAPIVKFWFYTISYLGYLLLFNYVI----EVRMDGWPSLQE 791
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                   3.4%;
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Score 137; DB 3; Length 1533; Pred. No. 0.0011; Mismatches 180; Indels 13
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Conservative

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Indels 138;

Gaps

22;

Gaps

17;

294

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GENERAL INFORMATION:
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APPLICANT: Blebm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
TITLE OF INVENTION: USES THEREOF
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
                                                              APPLICATION NUMBER: US/08/847,429 FILING DATE: 24-APR-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                              CLASSIFICATION: 530 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                      MEDIUM TYPE: rloppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: WordPerfect for Windows,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Carol Talkington Verser, ADDRESSEE: Heska Corporation STREET: 1825 Sharp Point Drive CITY: Fort Collins STATE: Colorado
                    NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459
                                                                                                                                                                               FILING DATE:
                                                                                                                                                                                                   APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WIVISYIVS----LALEKIREILMSEPGKL----SQKIKVW----LQEYWNITDLVAISTF
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  NUMBER:
                                                                                                                                                                                                   US/09/031,485
                                                                                                                                                                                                                                               Version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23, Appl Patent No. 582769
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent No.
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Best Local
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APPLICANT: Tang, Liang
APPLICANT: Blehm, E. S.
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TELEFAX: 97
INFORMATION FOR
                                  NAME: Verser, Carol Talkington REGISTRATION NUMBER: 37,459 REFERENCE/DOCKET NUMBER: HW-5 TELECOMMUNICATION INFORMATION: TELEPHONE: 970,493-7272
                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Wordberfect for Windows, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   174 YTPLHIAAKEGQEEVAAILMDHGTDKTLLTKKGFTPL-HLAAK-----YGNLPVAKSLLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            295 LTPLKLAAKEGKIEIFRHIL-------QREFSGLSHLSRKFTEWCYG--PVRVSLYD 342
                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                      APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                ZIP: 80525
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LLLDR---NADP--NARALN----GFTPLHIACKKNRIKIVELLLKYHAAIEATTESGLS 76
                                                                                                                                                                                                                                                                                                                                                                                                                                               Fort Collins
                                                                                                                                                                                                                                                                                                                                                                                                                          Colorado
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Application US/08847429A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Heska Corporation
1825 Sharp Point Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            303 amino acids
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                       970/484-9505
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Pred. No. 0.00031;
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; SEQUENCE CHARACTERISTICS:
; LENGTH: 303 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-847-429A-23
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                                                                                                                                           343 LAS-VDSCEENSVLEI-IAFH 361
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228 RGTPVDIEGKNQVTPLHVAAH 248
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